

OIKE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/825,751

DATE: 09/19/2001

TIME: 14:51:03

Input Set : A:\Cura-251.app

Output Set: N:\CRF3\09192001\I825751.raw

3 <110> APPLICANT: Vernet, Corine A. M.
 4 Burgess, Catherine E
 5 Fernandes, Elma R
 6 Taupier, Raymond J
 7 Quinn, Kerry E
 8 Spytek, Kimberly A
 9 Rastelli, Luca
 10 Herrman, John L
 12 <120> TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
 14 <130> FILE REFERENCE: 15966-750
 16 <140> CURRENT APPLICATION NUMBER: 09/825,751
 C--> 17 <141> CURRENT FILING DATE: 2001-09-07
 19 <150> PRIOR APPLICATION NUMBER: 60/194,314
 20 <151> PRIOR FILING DATE: 2000-04-03
 22 <150> PRIOR APPLICATION NUMBER: 60/225,693
 23 <151> PRIOR FILING DATE: 2000-08-16
 25 <160> NUMBER OF SEQ ID NOS: 85
 27 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply
 Corrected Diskette Needed

Sequence is
 unintelligible
 Gross Format Errors
 Contact Robert Wax
 at 703 308 4216

ERRORED SEQUENCES

1733 <210> SEQ ID NO: 22
 1734 <211> LENGTH: 20
 1735 <212> TYPE: DNA
 1736 <213> ORGANISM: Artificial Sequence
 1738 <220> FEATURE:
 1739 <223> OTHER INFORMATION: Description of Artificial Sequence:Forward Primer

E--> 1740 ag, 390

W--> 1742 <210> SEQ ID NO:

W--> 1742 <211> LENGTH:

W--> 1742 <212> TYPE:

W--> 1742 <213> ORGANISM:

1742 <400> SEQUENCE: 22

E--> 1743 accaatgtca tcggaggctt

1746 <210> SEQ ID NO: 23

1747 <211> LENGTH: 21

1748 <212> TYPE: DNA

1749 <213> ORGANISM: Artificial Sequence

1751 <220> FEATURE:

1752 <223> OTHER INFORMATION: Description of Artificial Sequence:Reverse Primer

E--> 1753 ag, 390

W--> 1755 <210> SEQ ID NO:

W--> 1755 <211> LENGTH:

W--> 1755 <212> TYPE:

W--> 1755 <213> ORGANISM:

1755 <400> SEQUENCE: 23

"ag" characters need to be
 at the end of line 1739

"ag" characters need to be at
 the end of line 1752

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E--> 1756 gatgtcctcg caggtcatca t 21
1759 <210> SEQ ID NO: 24
1760 <211> LENGTH: 23
1761 <212> TYPE: DNA
1762 <213> ORGANISM: Artificial Sequence
1764 <220> FEATURE:
1765 <223> OTHER INFORMATION: Description of Artificial Sequence:Probe Primer
E--> 1766 ag 290
W--> 1768 <210> SEQ ID NO:
W--> 1768 <211> LENGTH:
W--> 1768 <212> TYPE:
W--> 1768 <213> ORGANISM:
1768 <400> SEQUENCE: 24
E--> 1769 tcaaagccgt cagcacaggc aca
1787 <210> SEQ ID NO: 26
1788 <211> LENGTH: 21
1789 <212> TYPE: DNA
1790 <213> ORGANISM: Artificial Sequence
1792 <220> FEATURE:
1793 <223> OTHER INFORMATION: Description of Artificial Sequence:Forward Primer
E--> 1794 ag 271
W--> 1796 <210> SEQ ID NO:
W--> 1796 <211> LENGTH:
W--> 1796 <212> TYPE:
W--> 1796 <213> ORGANISM:
1796 <400> SEQUENCE: 26
E--> 1797 acctggacat agggcgtgtc t 21
1800 <210> SEQ ID NO: 27
1801 <211> LENGTH: 20
1802 <212> TYPE: DNA
1803 <213> ORGANISM: Artificial Sequence
1805 <220> FEATURE:
1806 <223> OTHER INFORMATION: Description of Artificial Sequence:Reverse Primer
E--> 1807 ag 271
W--> 1809 <210> SEQ ID NO:
W--> 1809 <211> LENGTH:
W--> 1809 <212> TYPE:
W--> 1809 <213> ORGANISM:
1809 <400> SEQUENCE: 27
E--> 1810 tcgatggaag tctccttgcc 20
1838 <210> SEQ ID NO: 30
1839 <211> LENGTH: 20
1840 <212> TYPE: DNA
1841 <213> ORGANISM: Artificial Sequence
1843 <220> FEATURE:
1844 <223> OTHER INFORMATION: Description of Artificial Sequence:Forward Primer
E--> 1845 ag 72
W--> 1847 <210> SEQ ID NO:
W--> 1847 <211> LENGTH:

"ag" characters need to be at end of line 1765.

Global Errors 23

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W--> 1847 <212> TYPE:
W--> 1847 <213> ORGANISM:
      1847 <400> SEQUENCE: 30
E--> 1848 cggaaagacc cagcagtgtt                20
      1851 <210> SEQ ID NO: 31
      1852 <211> LENGTH: 25
      1853 <212> TYPE: DNA
      1854 <213> ORGANISM: Artificial Sequence
      1856 <220> FEATURE:
      1857 <223> OTHER INFORMATION: Description of Artificial Sequence:Reverse Primer
E--> 1858 ag 72
W--> 1860 <210> SEQ ID NO:
W--> 1860 <211> LENGTH:
W--> 1860 <212> TYPE:
W--> 1860 <213> ORGANISM:
      1860 <400> SEQUENCE: 31
E--> 1861 atgatgtgaa cgagtgtgag tcctt          25
      1893 <210> SEQ ID NO: 34
      1894 <211> LENGTH: 22
      1895 <212> TYPE: DNA
      1896 <213> ORGANISM: Artificial Sequence
      1898 <220> FEATURE:
      1899 <223> OTHER INFORMATION: Description of Artificial Sequence:Forward Primer
E--> 1900 ag 248
W--> 1902 <210> SEQ ID NO:
W--> 1902 <211> LENGTH:
W--> 1902 <212> TYPE:
W--> 1902 <213> ORGANISM:
      1902 <400> SEQUENCE: 34
E--> 1903 tttccaacaa gatctgcaac ca            22
      1906 <210> SEQ ID NO: 35
      1907 <211> LENGTH: 17
      1908 <212> TYPE: DNA
      1909 <213> ORGANISM: Artificial Sequence
      1911 <220> FEATURE:
      1912 <223> OTHER INFORMATION: Description of Artificial Sequence:Reverse Primer
E--> 1913 ag 248
W--> 1915 <210> SEQ ID NO:
W--> 1915 <211> LENGTH:
W--> 1915 <212> TYPE:
W--> 1915 <213> ORGANISM:
      1915 <400> SEQUENCE: 35
E--> 1916 aggtagccccg cgcagag                17
      1947 <210> SEQ ID NO: 38
      1948 <211> LENGTH: 25
      1949 <212> TYPE: DNA
      1950 <213> ORGANISM: Artificial Sequence
      1952 <220> FEATURE:
      1953 <223> OTHER INFORMATION: Description of Artificial Sequence:Forward Primer

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E--> 1954 ag 287

W--> 1956 <210> SEQ ID NO:

W--> 1956 <211> LENGTH:

W--> 1956 <212> TYPE:

W--> 1956 <213> ORGANISM:

1956 <400> SEQUENCE: 38

E--> 1957 aactcagact gcaattgtga tgaaa

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L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1740 M:254 E: No. of Bases conflict, LENGTH:Input:390 Counted:2 SEQ:22
L:1740 M:112 C: (48) String data converted to lower case,
L:1740 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:22
L:1742 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1742 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1742 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1742 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1753 M:254 E: No. of Bases conflict, LENGTH:Input:390 Counted:2 SEQ:23
M:112 Repeated in SeqNo=23
L:1753 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:2 SEQ:23
L:1755 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1755 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1755 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1755 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1766 M:254 E: No. of Bases conflict, LENGTH:Input:390 Counted:2 SEQ:24
M:112 Repeated in SeqNo=24
L:1766 M:252 E: No. of Seq. differs, <211>LENGTH:Input:23 Found:2 SEQ:24
L:1768 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1768 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1768 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1768 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1794 M:254 E: No. of Bases conflict, LENGTH:Input:271 Counted:2 SEQ:26
M:112 Repeated in SeqNo=26
L:1794 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:2 SEQ:26
L:1796 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1796 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1796 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1796 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1807 M:254 E: No. of Bases conflict, LENGTH:Input:271 Counted:2 SEQ:27
M:112 Repeated in SeqNo=27
L:1807 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:27
L:1809 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1809 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1809 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1809 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1845 M:254 E: No. of Bases conflict, LENGTH:Input:72 Counted:2 SEQ:30
M:112 Repeated in SeqNo=30
L:1845 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:30
L:1847 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1847 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1847 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1847 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1858 M:254 E: No. of Bases conflict, LENGTH:Input:72 Counted:2 SEQ:31
M:112 Repeated in SeqNo=31
L:1858 M:252 E: No. of Seq. differs, <211>LENGTH:Input:25 Found:2 SEQ:31
L:1860 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1860 M:282 W: Numeric Field Identifier Missing, <211> is required.

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L:1860 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1860 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1900 M:254 E: No. of Bases conflict, LENGTH:Input:248 Counted:2 SEQ:34
M:112 Repeated in SeqNo=34
L:1900 M:252 E: No. of Seq. differs, <211>LENGTH:Input:22 Found:2 SEQ:34
L:1902 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1902 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1902 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1902 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1913 M:254 E: No. of Bases conflict, LENGTH:Input:248 Counted:2 SEQ:35
M:112 Repeated in SeqNo=35
L:1913 M:252 E: No. of Seq. differs, <211>LENGTH:Input:17 Found:2 SEQ:35
L:1915 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1915 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1915 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1915 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1954 M:254 E: No. of Bases conflict, LENGTH:Input:287 Counted:2 SEQ:38
M:112 Repeated in SeqNo=38
L:1954 M:252 E: No. of Seq. differs, <211>LENGTH:Input:25 Found:2 SEQ:38
L:1956 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1956 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1956 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1956 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:1967 M:254 E: No. of Bases conflict, LENGTH:Input:287 Counted:2 SEQ:39
M:112 Repeated in SeqNo=39
L:1967 M:252 E: No. of Seq. differs, <211>LENGTH:Input:28 Found:2 SEQ:39
L:1969 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1969 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1969 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1969 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:2007 M:254 E: No. of Bases conflict, LENGTH:Input:252 Counted:2 SEQ:42
M:112 Repeated in SeqNo=42
L:2007 M:252 E: No. of Seq. differs, <211>LENGTH:Input:19 Found:2 SEQ:42
L:2009 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:2009 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:2009 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:2009 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:2020 M:254 E: No. of Bases conflict, LENGTH:Input:252 Counted:2 SEQ:43
M:112 Repeated in SeqNo=43
L:2020 M:252 E: No. of Seq. differs, <211>LENGTH:Input:24 Found:2 SEQ:43
L:2022 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:2022 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:2077 M:254 E: No. of Bases conflict, LENGTH:Input:16 Counted:2 SEQ:46
M:112 Repeated in SeqNo=46
L:2077 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:46
L:2090 M:254 E: No. of Bases conflict, LENGTH:Input:16 Counted:2 SEQ:47
M:112 Repeated in SeqNo=47
L:2090 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:47
L:2103 M:254 E: No. of Bases conflict, LENGTH:Input:16 Counted:2 SEQ:48

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M:112 Repeated in SeqNo=48

L:2103 M:252 E: No. of Seq. differs, <211>LENGTH:Input:23 Found:2 SEQ:48

L:2133 M:254 E: No. of Bases conflict, LENGTH:Input:177 Counted:2 SEQ:50

M:112 Repeated in SeqNo=50

L:2133 M:252 E: No. of Seq. differs, <211>LENGTH:Input:18 Found:2 SEQ:50

L:2146 M:254 E: No. of Bases conflict, LENGTH:Input:177 Counted:2 SEQ:51

M:112 Repeated in SeqNo=51

L:2146 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:2 SEQ:51

L:2159 M:254 E: No. of Bases conflict, LENGTH:Input:177 Counted:2 SEQ:52

M:112 Repeated in SeqNo=52

L:2159 M:252 E: No. of Seq. differs, <211>LENGTH:Input:24 Found:2 SEQ:52

L:2236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57

L:2237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57

L:2247 M:254 E: No. of Bases conflict, LENGTH:Input:191 Counted:2 SEQ:58

L:2247 M:112 C: (48) String data converted to lower case,

L:2247 M:252 E: No. of Seq. differs, <211>LENGTH:Input:24 Found:2 SEQ:58

L:2260 M:254 E: No. of Bases conflict, LENGTH:Input:191 Counted:2 SEQ:59

M:112 Repeated in SeqNo=59

L:2260 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:2 SEQ:59

L:2291 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61